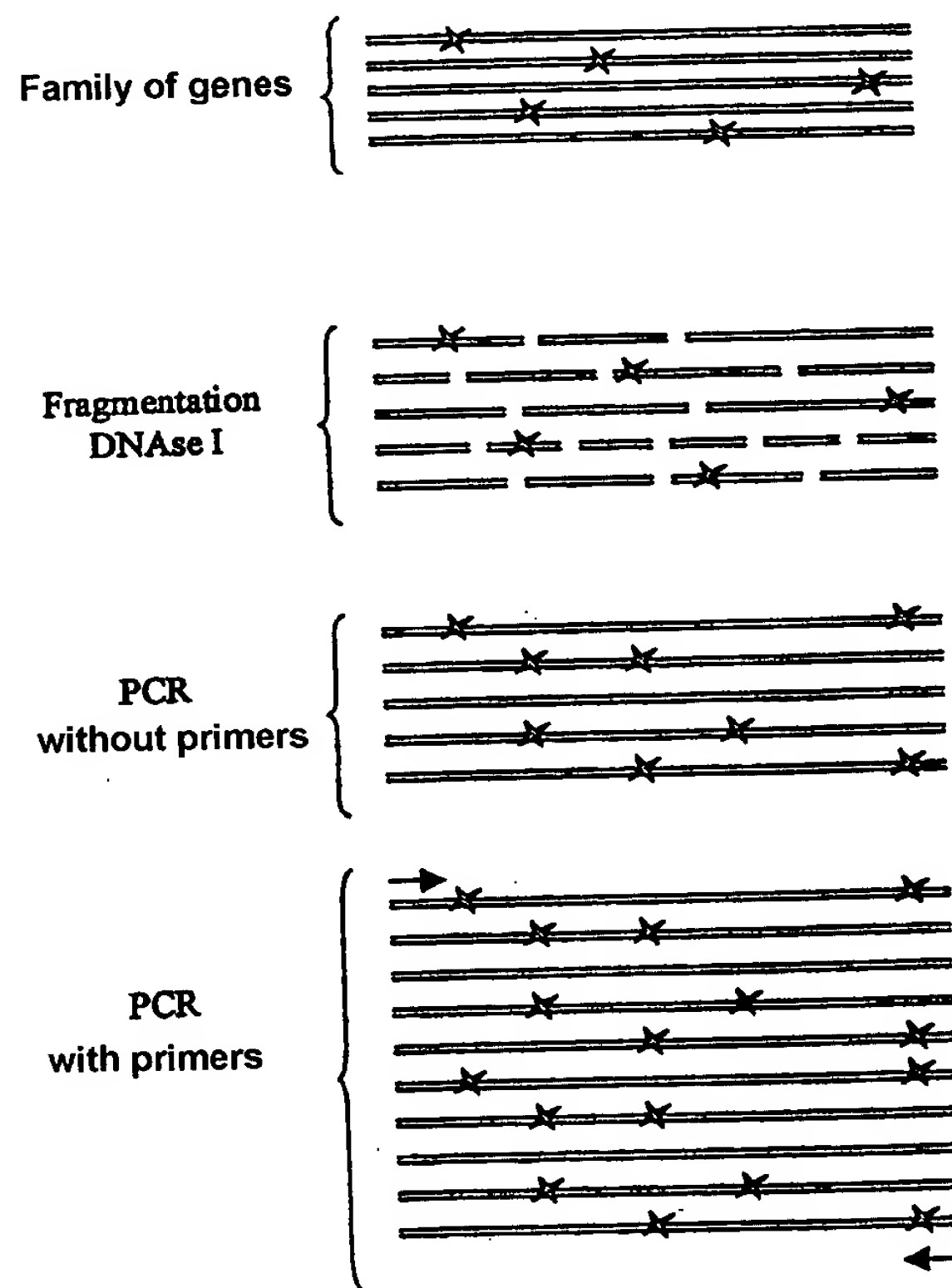


#4

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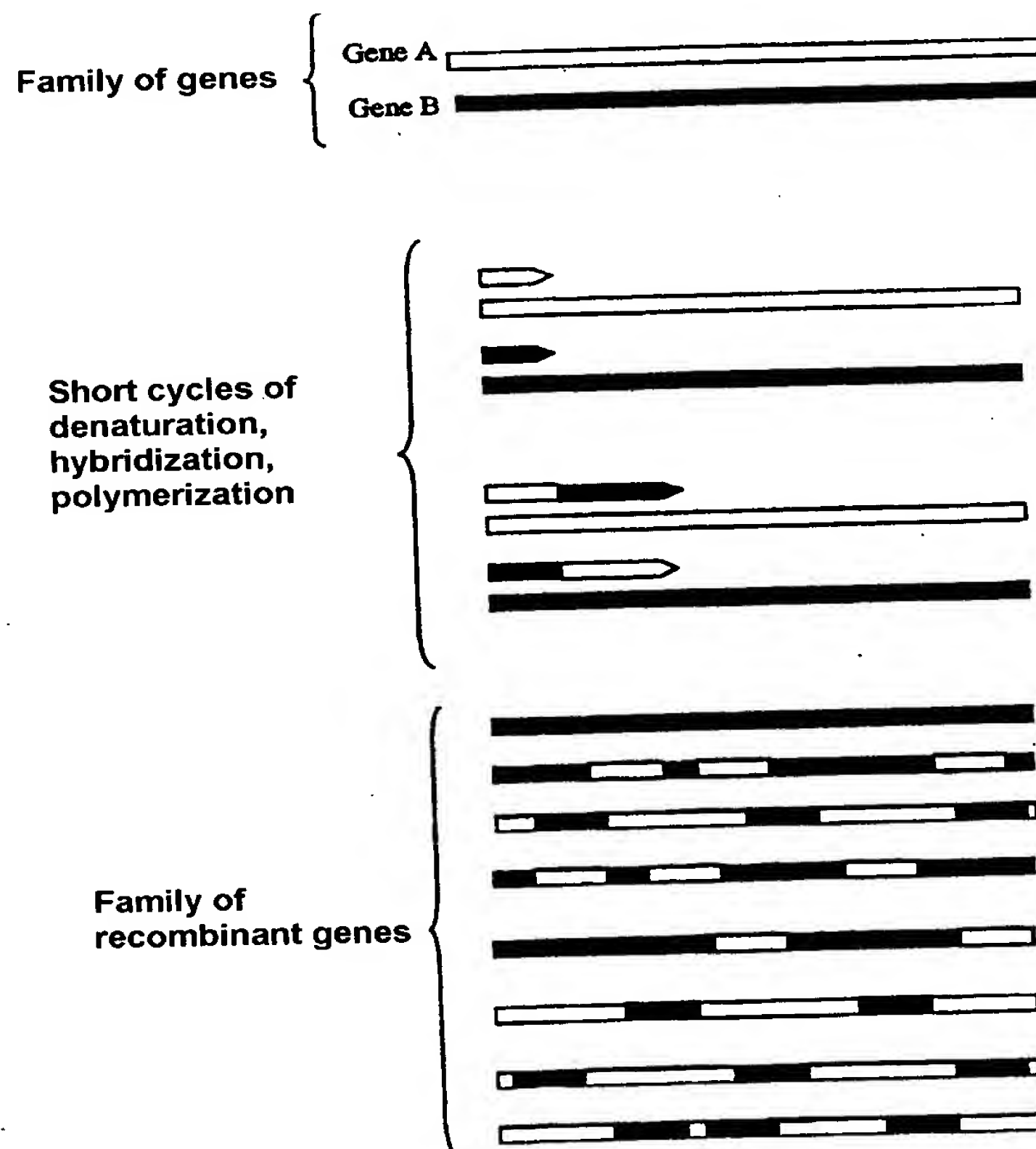
Figure 1 A



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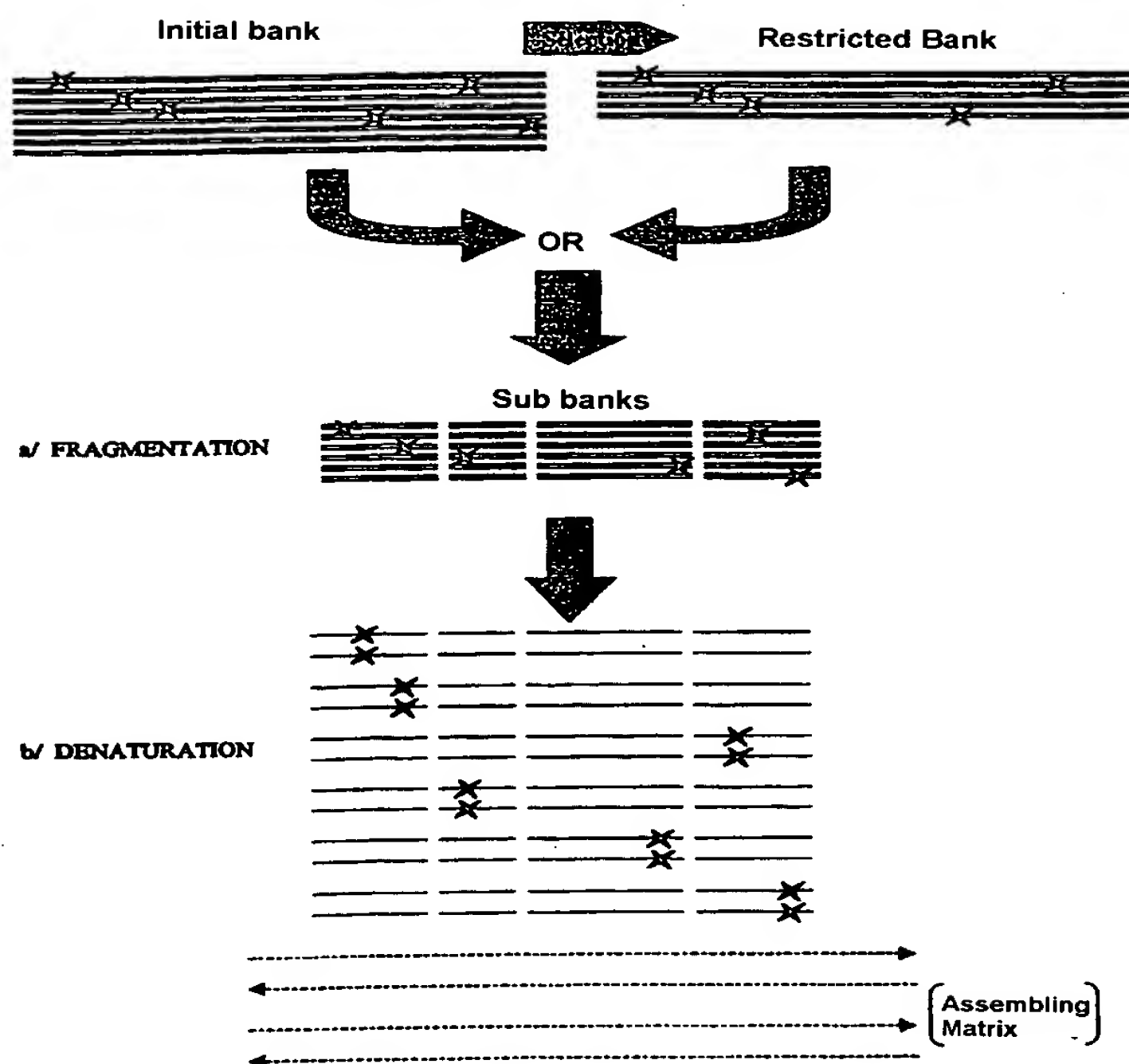
**Figure 1 B**

(Double-stranded process carried out, but illustrated here with a single strand)



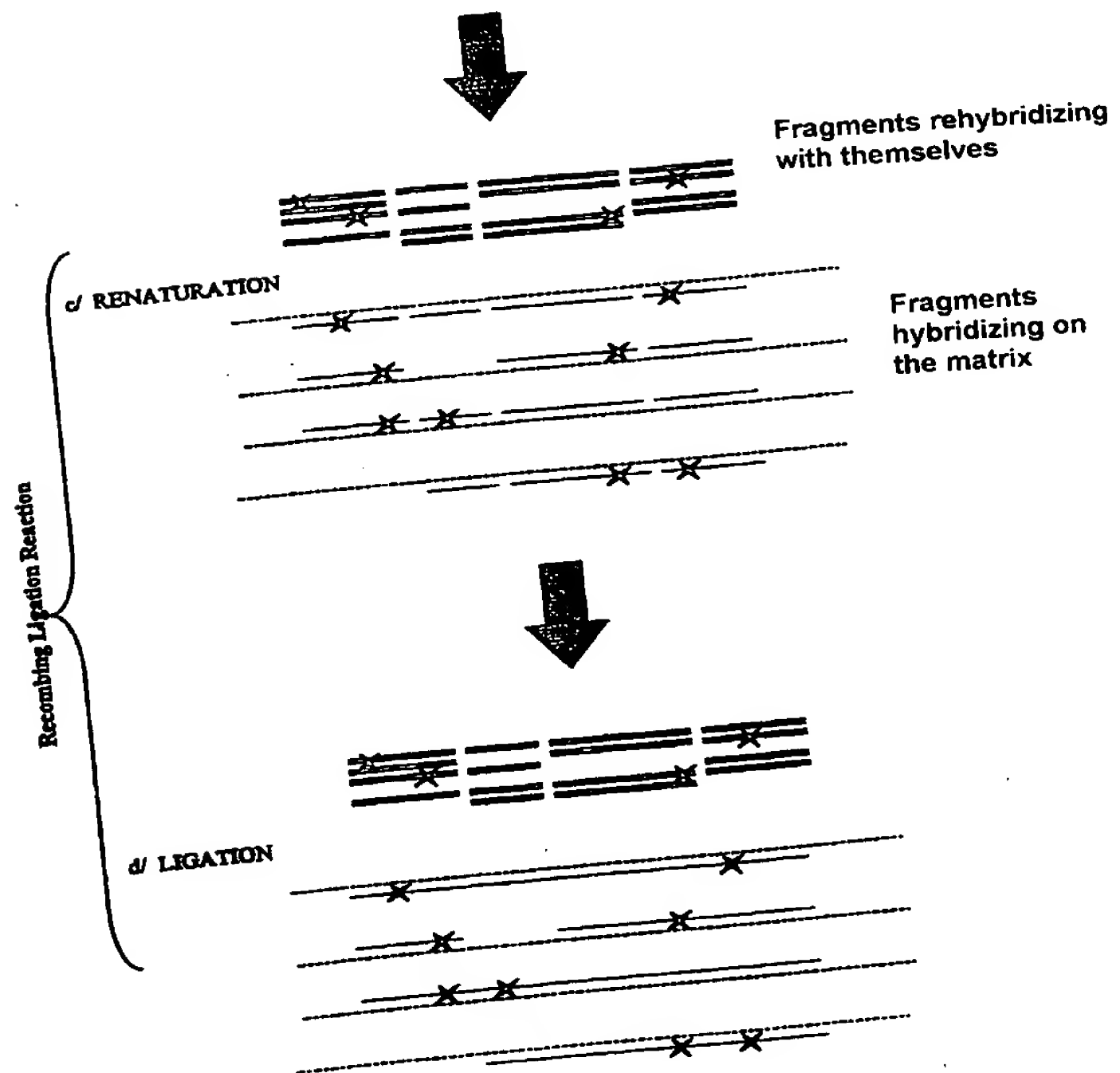
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Fig. 2



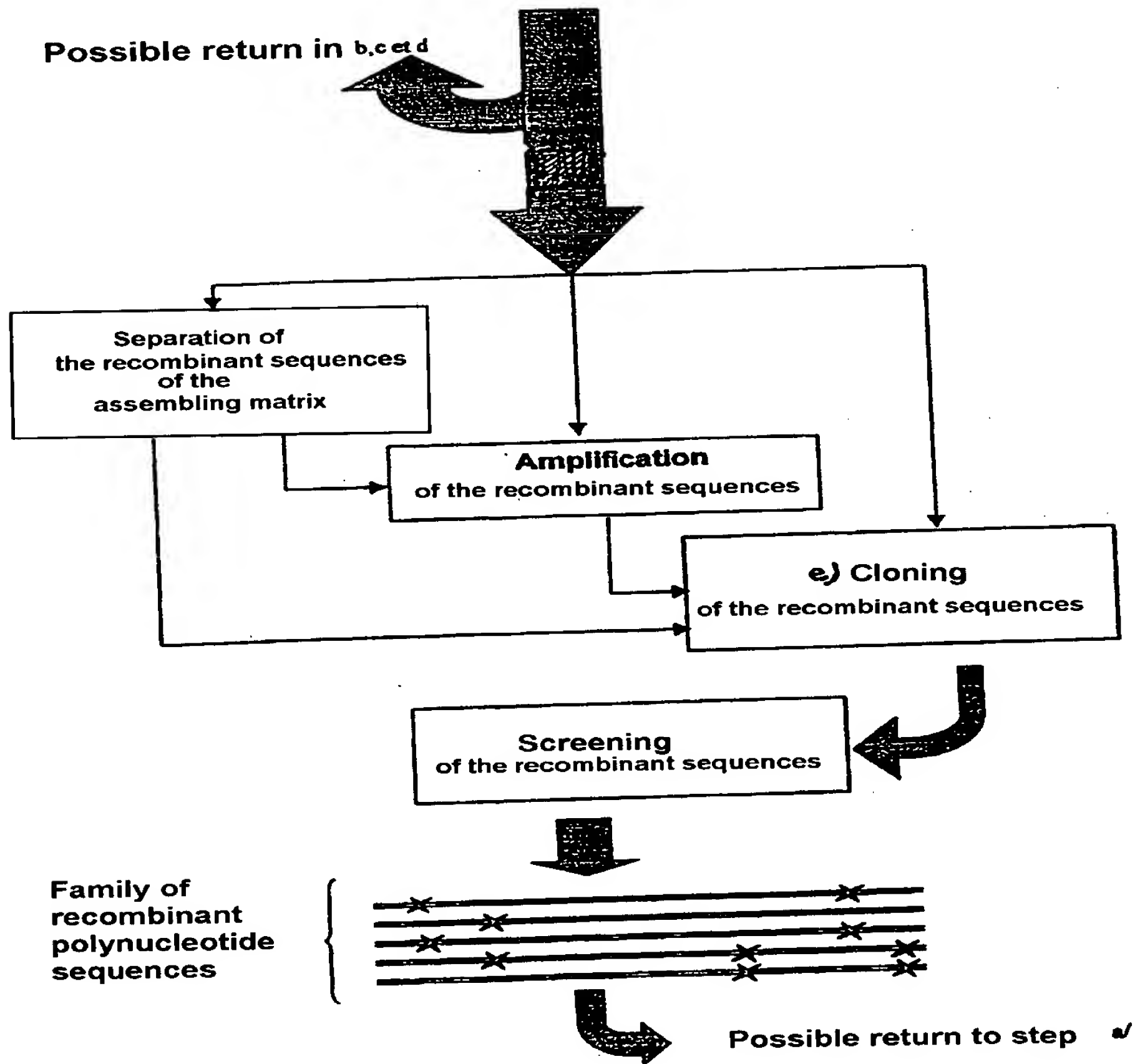
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Fig. 2- continued



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Fig. 2 continued



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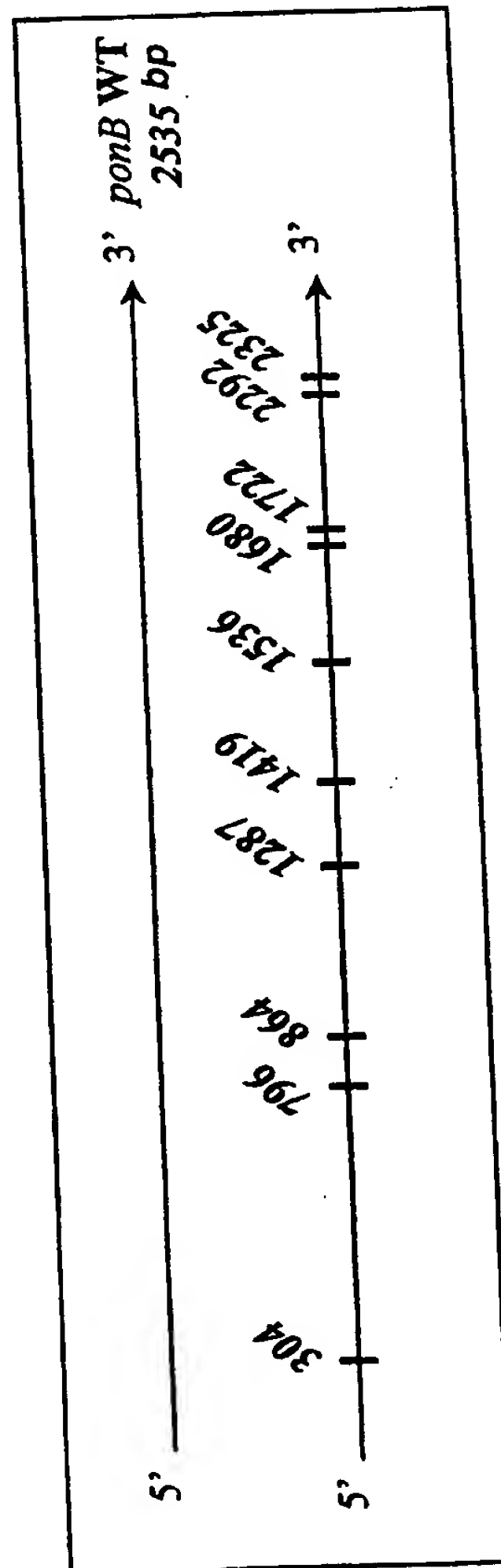


Figure 3: Position of the ten mutation zones (sites *Pvu II* and *Pst I*)

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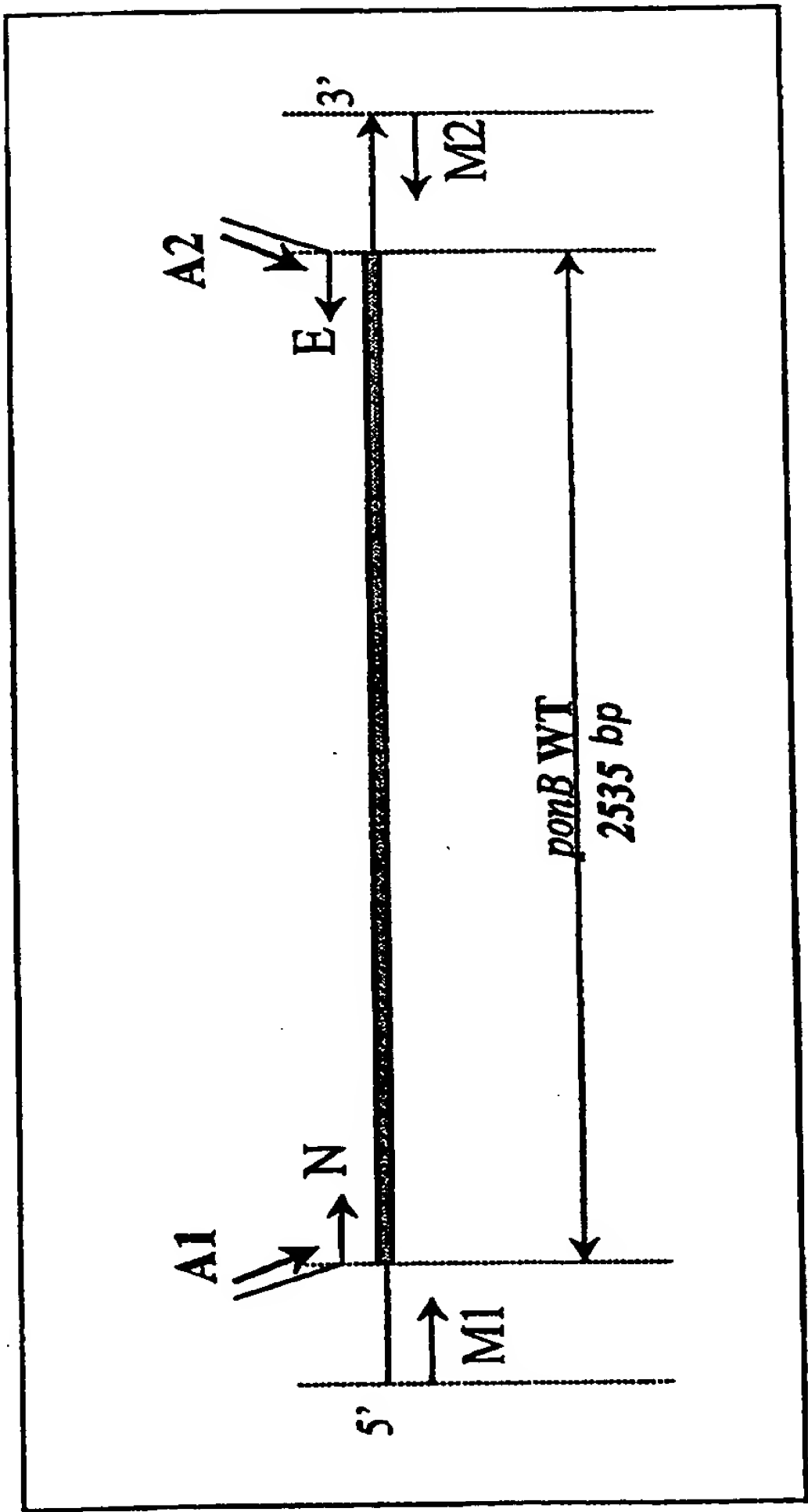
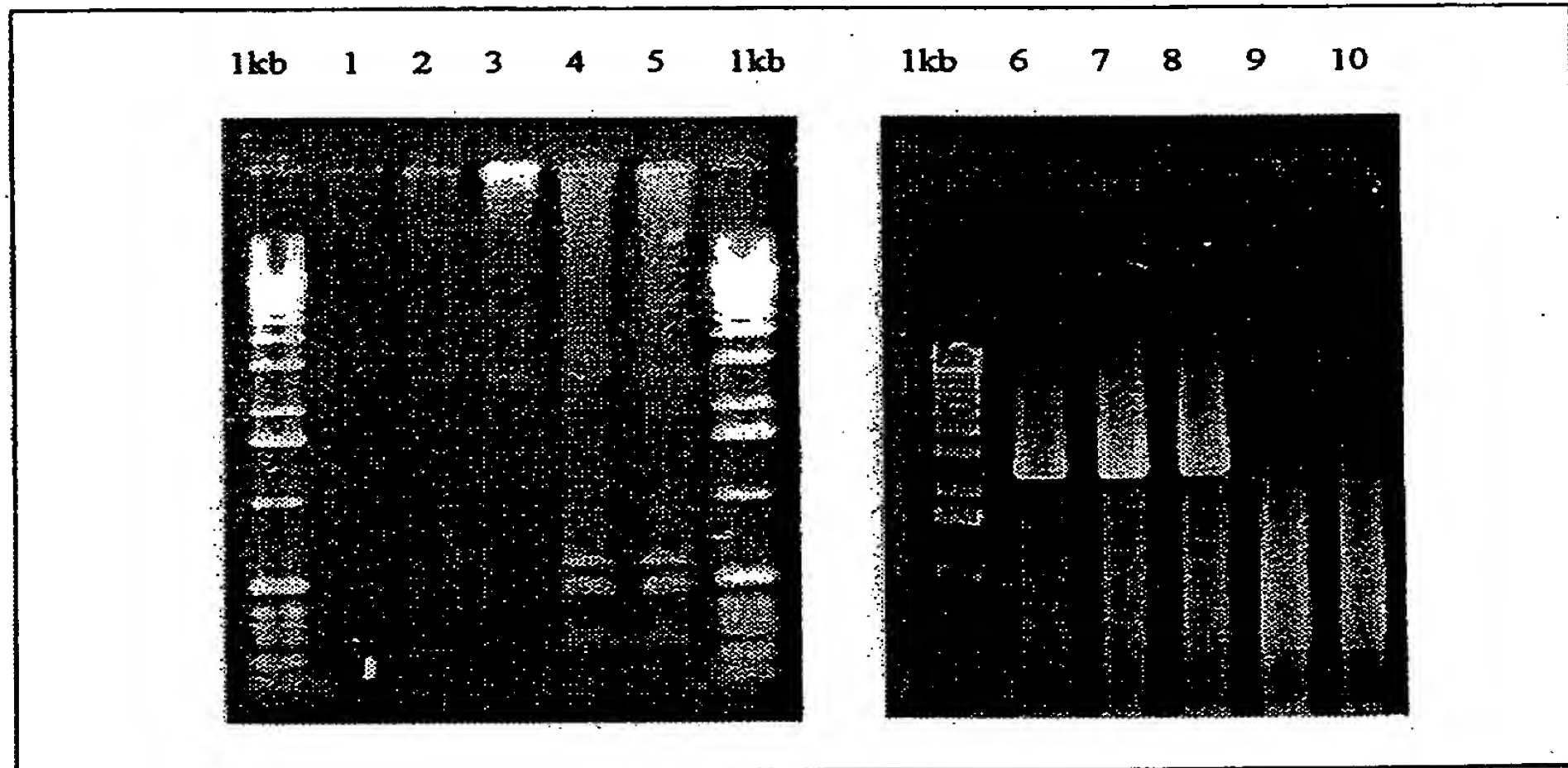


Figure 4 : Position of the primers used as compared to the sequence of the *ponB* gene

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**Fig. 5 : Migration of the RLR reactions and of the PCR amplifications of these reactions**

|                 |                       |                            |
|-----------------|-----------------------|----------------------------|
| <b>Tracks :</b> | <b>1/ RLR 1</b>       | <b>6/ PCR RLR 1</b>        |
|                 | <b>2/ RLR 2</b>       | <b>7/ PCR RLR 2</b>        |
|                 | <b>3/ RLR 3</b>       | <b>8/ PCR RLR 3</b>        |
|                 | <b>4/ RLR 4</b>       | <b>9/ PCR RLR 4</b>        |
|                 | <b>5/ RLR Control</b> | <b>10/ PCR RLR Control</b> |



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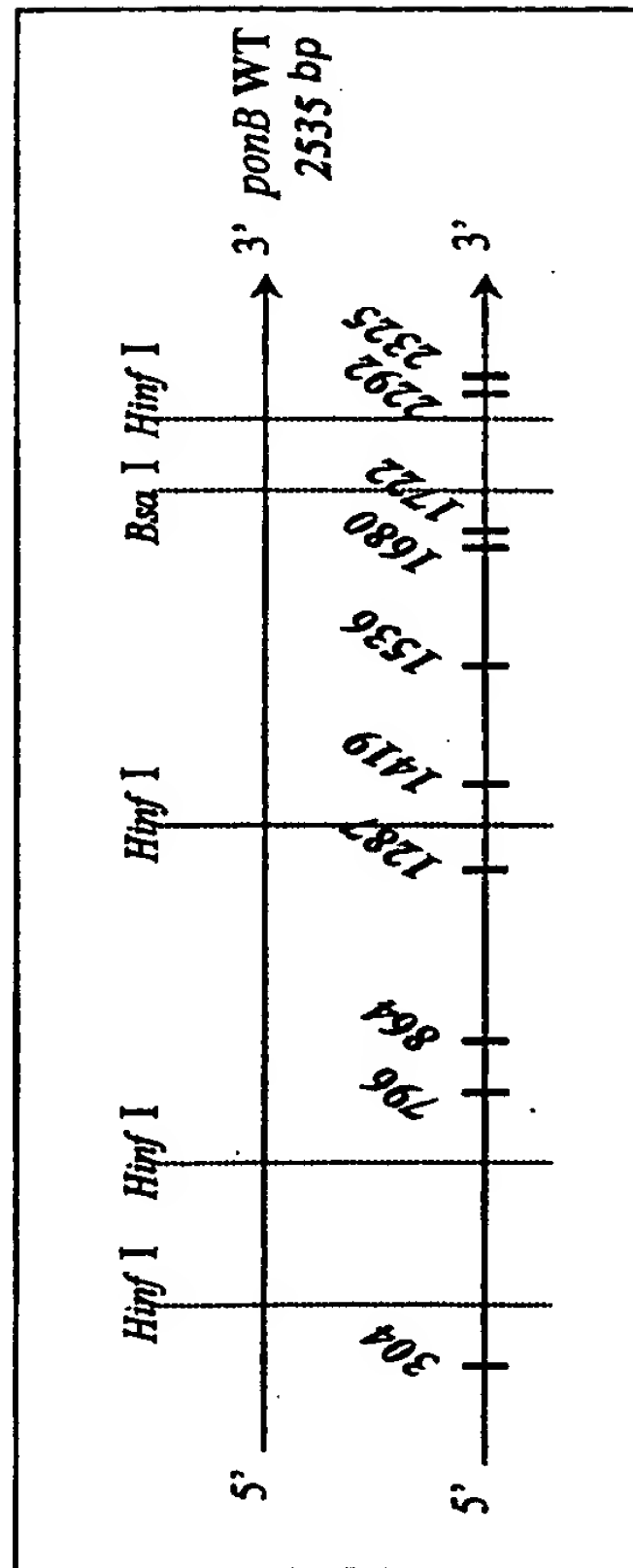


Figure 6 : Position of the mutations as compared to the restriction fragments